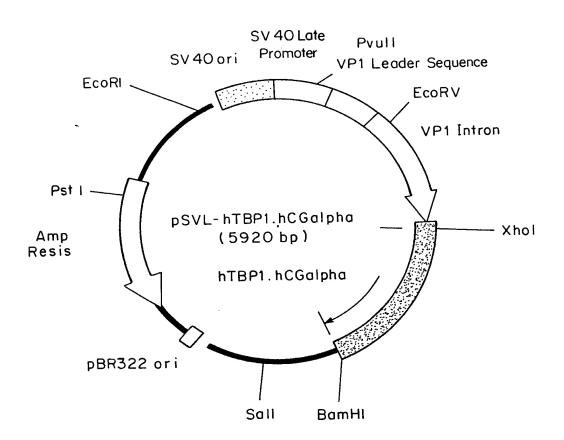


FIG. 1a(1)





F16.10(2)

hGH Signal Sequence

TCGAG

hGH Intron

CTT TGG CCC CTG TGC CTC CTG GGC G1y TTT GCT CTG CTC CTG TCC ACG CGG CGGCTCCCTCTGTTGCCCTCTGGTTTCTCCCCAGGC TCC

GGC AGT GCC GAT AGT GTG TGT CCC CAA GGA AAA TAT ATC CAC CCT CAA AAT AAT TCG ATT TGC TGT ACC AAG TGC CAC AAA AAA +20 Asp of Processed TBP1

ACC TAC TTG TAC AAT GAC TGT CCA GGC CCG GGG CAG GAT ACG GAC TGC AGG GAG TGT GAG AGC GGC TCC TTC ACC GCT TCA GAA AAC CAC CTC TAC TAC AST ASP CYS Pro Gly Pro Gly Gln Asp Thr Asp Cys Arg Glu Cys Glu Ser Gly Ser Phe Thr Ala Ser Glu Asn His Leu

Linker

G1yGGTPro CAG GAG AAA CAG AAC ACC GIG IGC ACC IGC CAI GCA GGI IIC III CIA AGA GAA AAC GAG IGI GIC IGI GCC GGI GCI GCC CCA Bgln Glu Lys Gln Asn Thr Val Cys Thr Cys His Ala Gly Phe Phe Leu Arg Glu Asn Glu Cys Val Ser Cys Ala Gly Ala Ala

TGC CCA GAA TGC ACG CTA CAG GAA AAC CCA TTC TTC TCC CAG CCG GGT GCC CCA ATA CTT CAG TGC ATG GGC TGC TGC TTC TCT AGA GCA Cys Pro Glu Cys Thr Leu Gln Glu Asn Pro Phe Phe Ser Gln Pro Gly Ala Pro Ile Leu Gln Cys Met Gly Cys Cys Phe Ser Arg Ala +7 Cys of hCG alpha

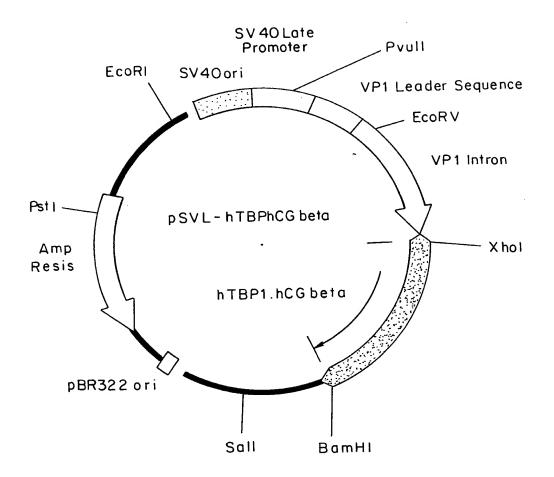
TCC AAG AAG ACG ATG TTG GTC CAA AAG AAC GTC ACT TCA GAG TCC ACT TGC TGT GTA GCT AAA TCA TAT AAC AGG GTC Ser Lys Lys Thr Met Leu Val Gln Lys Asn Val Thr Ser Glu Ser Thr Cys Cys Val Ala Lys Ser Tyr Asn Arg Val CCC ACT CCA CTA AGG
Pro Thr Pro Leu Arg

Ö ACA GTA ATG GGG GGT TTC AAA GTG GAG AAC CAC ACG GCG TGC CAC TGC AGT ACT TGT TAT TAT CAC AAA TCT TAA C

Bam HI



FIG. 1b(1)





F1G.16/21

hGH Signal Sequence

XhoI

hGH Intron

crcGag ATG GCT ACA Met Ala Thr TGC GGC CTG CTC Gly Leu Leu TTT CTG CTC CTG GCT Leu Leu Leu Ala TCC CGG ACG TCC Ser Arg Thr Ser CTCTTGCTCTCCGGCTCCCTCTGTTGCCCTCTGGTTTCTCCCCAGGC

+20 Asp of Processed TBP1

ACC TGI TGC ATT 700 AAT AAT CAA CCI CAC GAT AGT GTG TGT CCC CAA GGA AAA TAT ATC CCC TGG CTT CAA GAG GGC AGT GCC Pro Trp Leu Gln Glu Gly Ser Ala

Asp Ser Val Cys Pro Gln Gly Lys Tyr Ile His Pro Gln Asn Asn Ser Ile Cys Cys Thr

AAG TGC CAC AAA GGA ACC TAC TTG TAC AAT GAC TGT CCA GGC CCG GGG CAG GAT ACG GAC TGC AGG GAG TGT GAG AGC GGC TCC TTC ACC LYS CYS His Lys Gly Thr Tyr Leu Tyr Asn Asp Cys Pro Gly Gln Asp Thr Asp Cys Arg Glu Cys Glu Ser Gly Ser Phe Thr GCT TCA GAA AAC CAC CTC AGA CAC TGC CTC AGC TGC TCC AAA TGC CGA AAG GAA ATG GGT CAG GTG Ala Ser Glu Asn His Leu Arg His Cys Leu Ser Cys Ser Lys Cys Arg Lys Glu Met Gly Gln Val

GAG ATC TCT TGC ACA GTG GAC Glu Ile Ser Ser Cys Thr Val Asp

TGC CTC Cys Leu CGG GAC ACC GTG TGT GGC TGC AGG AAG AAC CAG TAC CGG CAT TAT TGG AGT GAA AAC CTT TTC CAG TGC TTC AAT TGC AGC CTC Arg Asp Thr Val Cys Gly Cys Arg Lys Asn Gln Tyr Arg His Tyr Trp Ser Glu Asn Leu Phe Gln Cys Phe Asn Cys Ser Leu GGG ACC GTG CAC CTC TCC TGC CAG GAG AAA CAG AAC ACC GTG TGC ACC TGC CAT GCA GGT TTC TTT CTA AGA GAA AAT GAG TGT GTC Gly Thr Val His Leu Ser Cys Gln Glu Lys Gln Asn Thr Val Cys Thr Cys His Ala Gly Phe Phe Leu Arg Glu Asn Glu Cys Val +7 Pro of hCG beta AAT Asn

Ser Cys Ala Gly Ala Gly Pro Arg Cys Arg Pro Ile Asn Ala Thr Leu Ala Val Glu Lys Glu Gly Cys Pro Val Cys Ile Thr Val TGC ATC ACC CCC GAG AAG GAG GGC TGC GCT GTG CIG TGT GCT GGT GGT CCA CGG TGC CGC CCC ATC AAT GCC ACC TCC

TGC AAC TAC Cys Asn Tyr AAC ACC AAC ATC TGT GCC GGC TAC TGC CCC ACC ATG ACC CGC GTG CTG CAG GGG GTC CTG CCG GCC CTG CCT CAG GTG GTG ASn Thr Thr Ile Cys Ala Gly Tyr Cys Pro Thr Met Thr Arg Val Leu Gln Gly Val Leu Pro Ala Leu Pro Gln Val Val Val

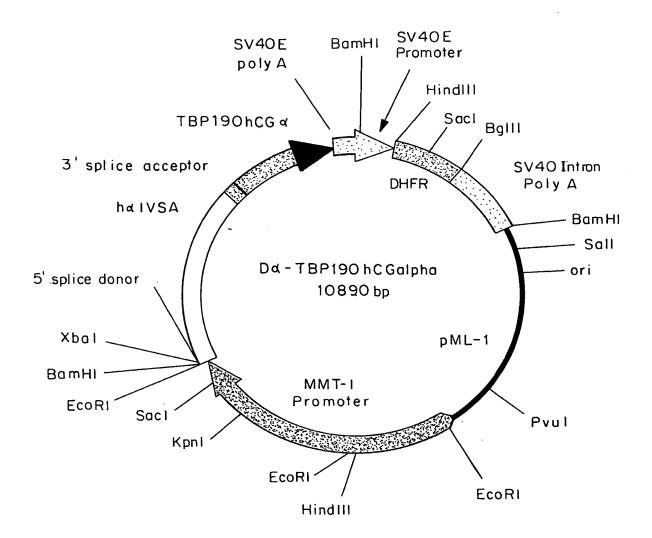
CGC CGC AGC ACC ACT GAC TGC GGG GGT CCC AAG GAC CAC CCC TTG ACC TGT GAT GAC CCC CGC TTC CAG GAC TCC TCT ANG ANG ANG Ser Thr Thr Asp Cys Gly Gly Pro Lys Asp His Pro Leu Thr Cys Asp Asp Pro Ang Phe Gln Asp Ser Ser CGC GAT GTG CGC TTC GAG TCC ATC CGG CTC CCT GGC TGC CGC CGC GGC GTG AAC CCC GTG GTC TCC TAC GCC GTG GCT CTC AGC TGT CAA Arg Asp Val Arg Phe Glu Ser Ile Arg Leu Pro Gly Cys Pro Arg Gly Val Asn Pro Val Val Ser Tyr Ala Val Ala Leu Ser Cys Gln TGC CGC CGC A TGT GCA CTC T Cys Ala Leu C

TCA AAG GCC CCT CCC CCC AGC CTT CCA AGC CCA TCC CGA CTC CCG GGG CCC TCG GAC ACC CCG ATC CTC CCA CAA TAA Ser Lys Ala Pro Pro Pro Ser Leu Pro Ser Arg Leu Pro Gly Pro Ser Asp Thr Pro Ile Leu Pro Gln ***

Bam HI



FIG. 2a(1)





F16.2a(2)

hGH Signal Sequence

GTAAGCGCCCCTAAAATCCCTTTGGGCACAATGTGTCCTGAGGGAAGAGGCAGCGACCTGTAGATGGGACGGGGGCACTAACCCTCAGGTTTGGGGTTTCT TCGAG ATG GCT ACA G

▶Met Ala Thr

CGGCTCCCTCTGTTGCCCTCTGGTTTCTCCCCAGGC TCC CGG ACG TCC CTG CTC CTG GCT TTT GGC CTG CTC TGC CTG CCT TGG CTT

+20 Asp of processed TBP1

CAA GAG GGC AGT GCC GAT AGT GTG TGT CCC CAA GGA AAA TAT ATC CAC CCT CAA AAT AAT TCG ATT TGC TGT ACC AAG TGC CAC AAA GGA GIN GIN GIU GIY Ser Ala Asp Ser Val Cys Pro Gin Gly Lys Tyr Ile His Pro Gin Asn Asn Ser Ile Cys Cys Thr Lys Cys His Lys Gly

ACC TAC TTG TAC AAT GAC TGT CCA GGC CCG GGG CAG GAT ACG GAC TGC AGG GAG TGT GAG AGC GGC TCC TTC ACC GCT TCA GAA AAC CAC CTC

CAG GAG AAA CAG AAC ACC GTG TGC ACC TGC CAT GCA GGT TTC TTT CTA AGA GAA AAC GAG TGT GTC TCC TGT AGT AAC TGT AAG AAA AGC CTG
GIn Glu Lys Gln Asn Thr Val Cys Thr Cys His Ala Gly Phe Phe Leu Arg Glu Asn Glú Cys Val Ser Asn Cys Lys Lys Ser Leu Linker

GCC CCA GGT TGC CCA Ala Pro Gly Cys Pro +7 Cys of hCG alpha GAG TGC ACG AAG TTG TGC CTA CCC CAG ATT GAG AAT GTT AAG GGC ACT GAG GAC TCA GGC ACC ACA GCC GGT GCT

GAA TGC ACG CTA CAG GAA AAC CCA TTC TCC CAG CCG GGT GCC CCA ATA CTT CAG TGC ATG GGC TGC TGC TTC TCT AGA GCA TAT CCC ACT

CCA CTA AGG TCC AAG AAG ACG ATG TTG GTC CAA AAG AAC GTC ACC TCA GAG TCC ACT TGC TGT GTA GCT AAA TCA TAT AAC AGG GTC ACA GTA

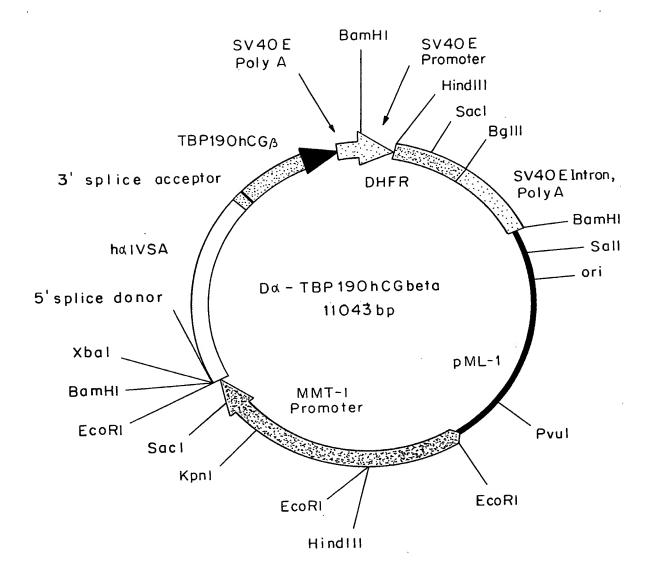
Pro Leu Arg Ser Lys Lys Thr Met Leu Val Gln Lys Asn Val Thr Ser Glu Ser Thr Cys Cys Val Ala Lys Ser Tyr Asn Arg Val Thr Val

ATG GGG GGT TTC AAA GTG GAG AAC CAC ACG GCG TGC CAC TGC AGT ACT TGT TAT TAT CAC AAA TCT TAA GGATCCCTCGAG

Wet Gly Gly Phe Lys Val Glu Asn His Thr Ala Cys His Cys Ser Thr Cys Tyr His Lys Ser ***



FIG. 2b(1)





F16. 26(2)

CTCGAG ATG GCT ACA G GTAAGCGCCCCTAAAATCCCTTTGGGCACAATGTGTCCTGAGGGGAGAAGCAGCGACCTGTAGATGGGGACGGGGGCACTAACCCTCAGGTTTGGG

CTG CTC CTG 66c G1y TTT GCT CTG CTC CTG TCC C TCC CGG ACG ▶ Ser Arg Thr CTCTTGCTCTCCGGCTCCCTCTGTTGCCCTCTGGTTTCTCCCCAGG C TCC

GAT AGT GTG TGT CCC CAA GGA AAA TAT ATC CAC CCT CAA AAT AAT TCG ATT TGC TGT ACC +20 Asp of Processed TBP1

Asp Ser Val Cys Pro Gln Gly Lys Tyr Ile His Pro Gln Asn Asn Ser Ile Cys Cys Thr TGG CTT CAA GAG GGC AGT GCC Trp Leu Gln Glu Gly Ser Ala

Pro '

TTC ACC Phe Thr AAG TGC CAC AAA GGA ACC TAC TTG TAC AAT GAC TGT CCA GGC CCG GGG CAG GAT ACG GAC TGC AGG GAG TGT GAG AGC GGC TCC Lys Cys His Lys Gly Thr Tyr Leu Tyr Asn Asp Cys Pro Gly Pro Gly Gln Asp Thr Asp Cys Arg Glu Cys Glu Ser Gly Ser

GCT TCA GAA AAC CAC CTC AGA CAC TGC CTC AGC TGC TCC AAA TGC CGA AAG GAA ATG GGT CAG GTG GAG ATC TCT TGC ACA GTG GAC

AAT GGG ACC GTG CAC TC TCC TGC CAG GAG AAA CAG AAC ACC GTG TGC ACC TGC CAT GCA GGT TTC TTT CTA AGA GAA AAC GAG TGT GTC

TCC TGT AGT AAC TGT AAG AAA AGC CTG GAG TGC ACG AAG TTG TGC CTA CCC CAG ATT GAG AAT GTT AAG GGC ACT GAG GAC TCA GGC ACC

Ser Cys Ser Asn Cys Lys Lys Ser Leu Glu Cys Thr Lys Leu Cys Leu Pro Gln Ile Glu Asn Val Lys Gly Thr Glu Asp Ser Gly Thr +7 Pro of beta ACA GCT GGT GCT GGT CCA CGG TGC CGC CCC ATC AAT GCC ACC CTG GCT GTG GAG AAG GAG GGC TGC CCC GTG TGC ATC ACC GTC AAC Thr Ala Gly Ala Gly Pro Arg Cys Arg Pro Ile Asn Ala Thr Leu Ala Val Glu Lys Glu Gly Cys Pro Val Cys Ile Thr Val Asn ACC ACC ATC TGT GCC GGC TAC TGC CCC ACC ATG ACC CGC GTG CTG CAG GGG GTC CTG CCG GCC CTG CCT CAG GTG GTG TGC AAC TAC CGC

The The Ile Cys Ala Gly Tyr Cys Pro The Met The Arg Val Leu Gln Gly Val Leu Pro Ala Leu Pro Gln Val Val Cys Asn Tyr Arg

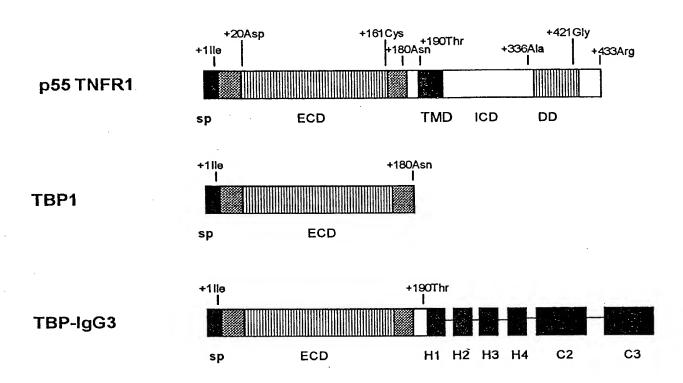
Asp Val Arg Phe Glu Ser Ile Arg Leu Pro Gly Cys Pro Arg Gly Val Asn Pro Val Val Ser Tyr Ala Val Ala Leu Ser Cys Gln Cys GAT GTG CGC TTC GAG TCC ATC CGG CTC CCT GGC TGC CCG CGC GGC GTG AAC CCC GTG GTC TCC TAC GCC GTG GCT CTC AGC TGT CAA TGT

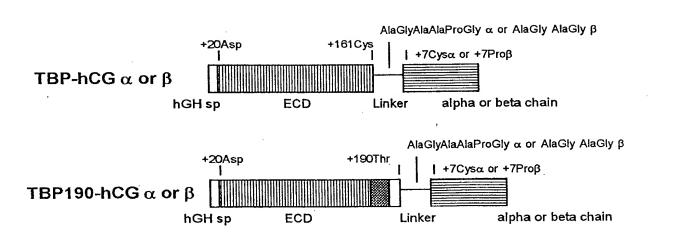
GCA CIC TGC CGC CGC AGC ACT GAC TGC GGG GGT CCC AAG GAC CAC CCC TTG ACC TGT GAT'GAC CCC CGC TTC CAG GAC TCC TCT TCC Ala Leu Cys Arg Arg Ser Thr Thr Asp Cys Gly Gly Pro Lys Asp His Pro Leu Thr Cys Asp Asp Pro Arg Phe Gln Asp Ser

GGATCCCTCGAG TCA AAG GCC CCT CCC CCC AGC CTT CCA AGC CCA TCC CGA CTC CCG GGG CCC TCG GAC ACC CCG ATC CTC CCA CAA TAA Ser Lys Ala Pro Pro Pro Ser Leu Pro Ser Pro Ser Arg Leu Pro Gly Pro Ser Asp Thr Pro Ile Leu Pro Gln ***



FIG. 3
p55 TNFR1, TBP1 and TBP1 FUSION CONSTRUCTS





CELLS + COS7 MOCK TRANSFECTANT MEDIA+2.5 ng/ml TNF4 -▼- CELLS + TBP-hCG(20-190) COS7 MED+2.5 ng/ml TNF --- 105 CELLS / WELL + 2.5 ng/ml TNF4 + TBP MONOMER TBP MONOMER CELLS ALONE CELLS + 2.5ng/ml TNF4 (NO TBP) TBP-hCG (20 - 190) CELLS + TNF (+/- MOCK MEDIA) CELLS ALONE 4 .0 – 1.0.1 2.0-4.57 3.5 3.0. 2.5

F16.4

ng/mi TBP EQUIVALENTS(R&D SYSTEM ELISA)

000

§

9

0.1

0.0

0.5

mn 078 **©** 0 0

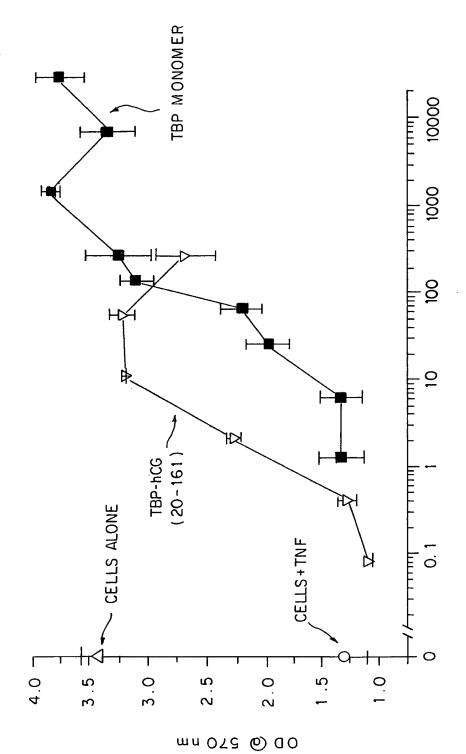




F16.5

Δ CELLS ALONE C CELLS + 2.5 ng/ml TNFα (NO TBP)

-7- CELLS+ PURIFIED TBP-hCG(20-161)



ng/ml TBP EQUIVALENTS (R&D SYSTEM ELISA)

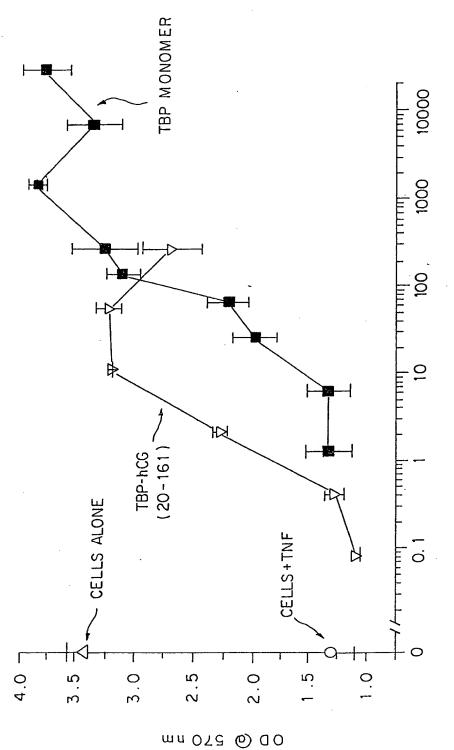
-E-105 CELLS / WELL +2.5 ng/ml TNFA + TBP MONOMER

A CELLS ALONE

F1G. 6

O CELLS + 2.5 ng/ml TNF4 (NO TBP)

-7- CELLS + PURIFIED TBP-hCG (20-161)



ng/ml TBP EQUIVALENTS (R&D SYSTEM ELISA)